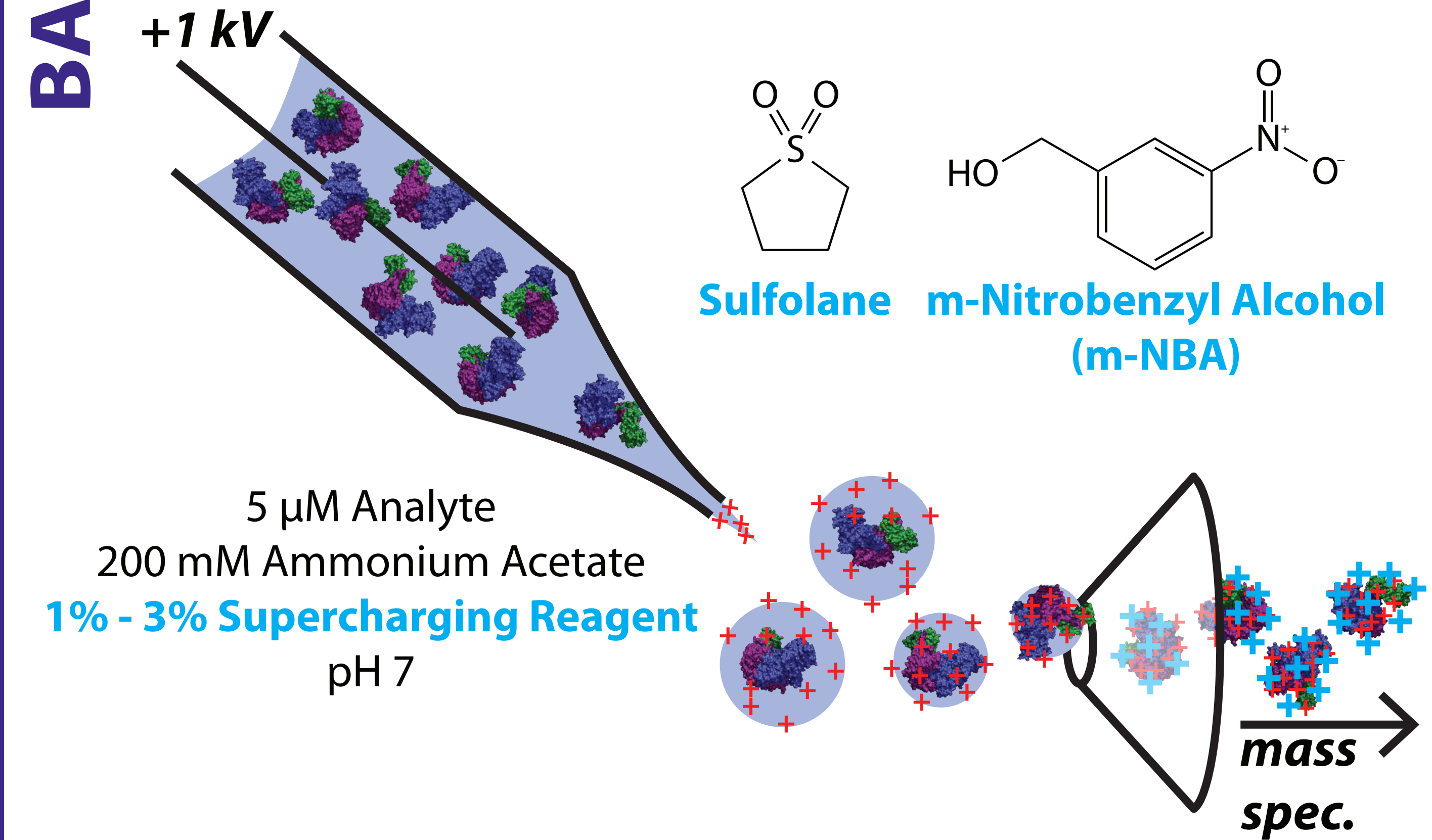


# Supercharging of Native-Like Proteins and Protein Complexes: Effects of m-Nitrobenzyl Alcohol versus Sulfolane

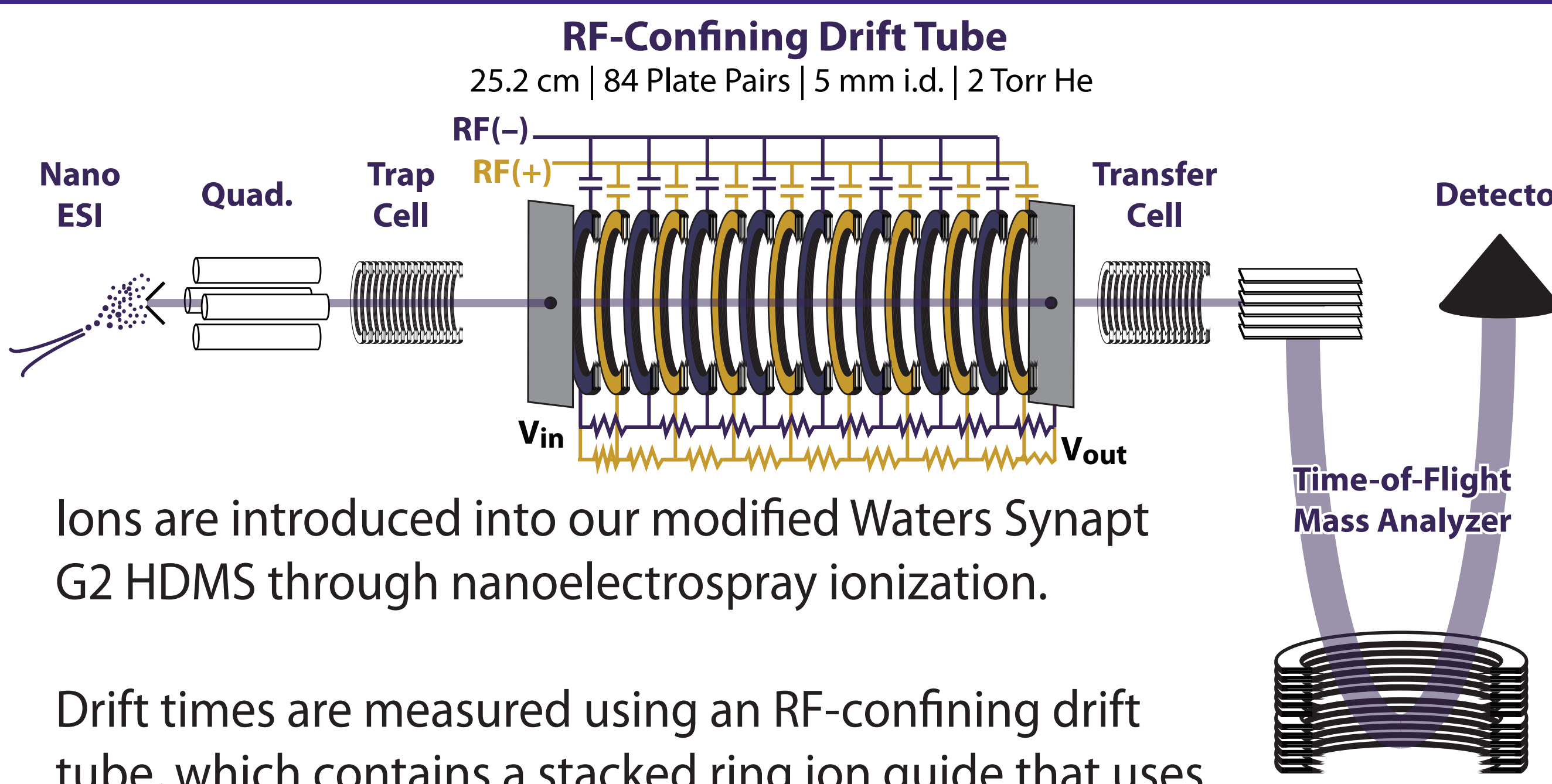
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**BACKGROUND** Electrospray ionization enables the transport of intact protein and protein complex ions from solution to the gas phase. Typically, electrospray ionization from ammonium acetate buffer solutions produces native mass spectra containing narrow charge-state distributions with low z-values. Certain organic molecules, however, can shift the charge-state distribution of electrosprayed proteins to higher z-values that are not usually accessible under native conditions. Here, we use sulfolane and m-nitrobenzyl alcohol (m-NBA) to **supercharge** proteins and protein complexes.

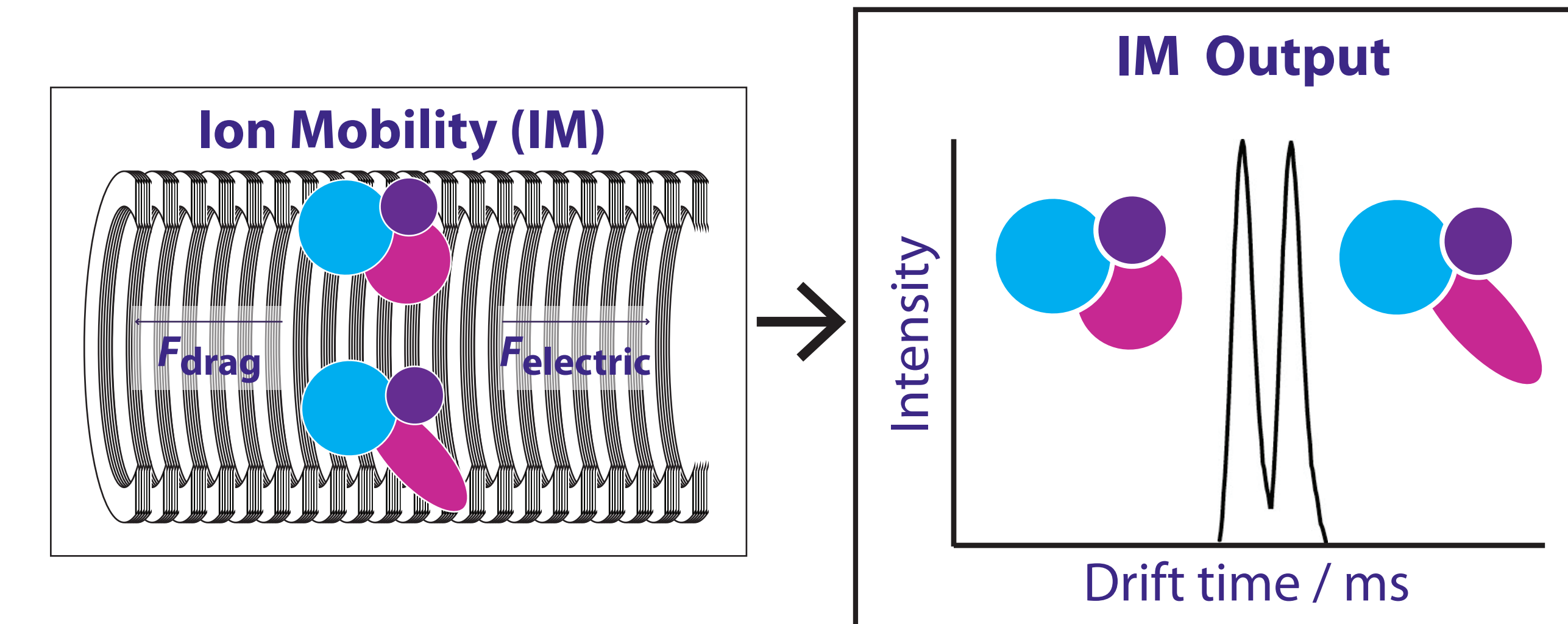


What are the effects of supercharging on the charge states and structures of proteins and protein complexes?

**INSTRUMENTATION & ION MOBILITY** Ions are introduced into our modified Waters Synapt G2 HDMS through nano-electrospray ionization.



Drift times are measured using an RF-confining drift tube, which contains a stacked ring ion guide that uses alternating RF potentials to radially confine ions and a superimposed DC potential to establish a uniform electric field along the axis of transmission.



Ion mobility separates ions based on their mobility, which depends on structure and charge. Ions with larger structures will experience more collisions with the neutral background gas and thus have longer drift times than smaller, more compact ions. Different ion conformations result in different drift times.

**SUPERCHARGING**

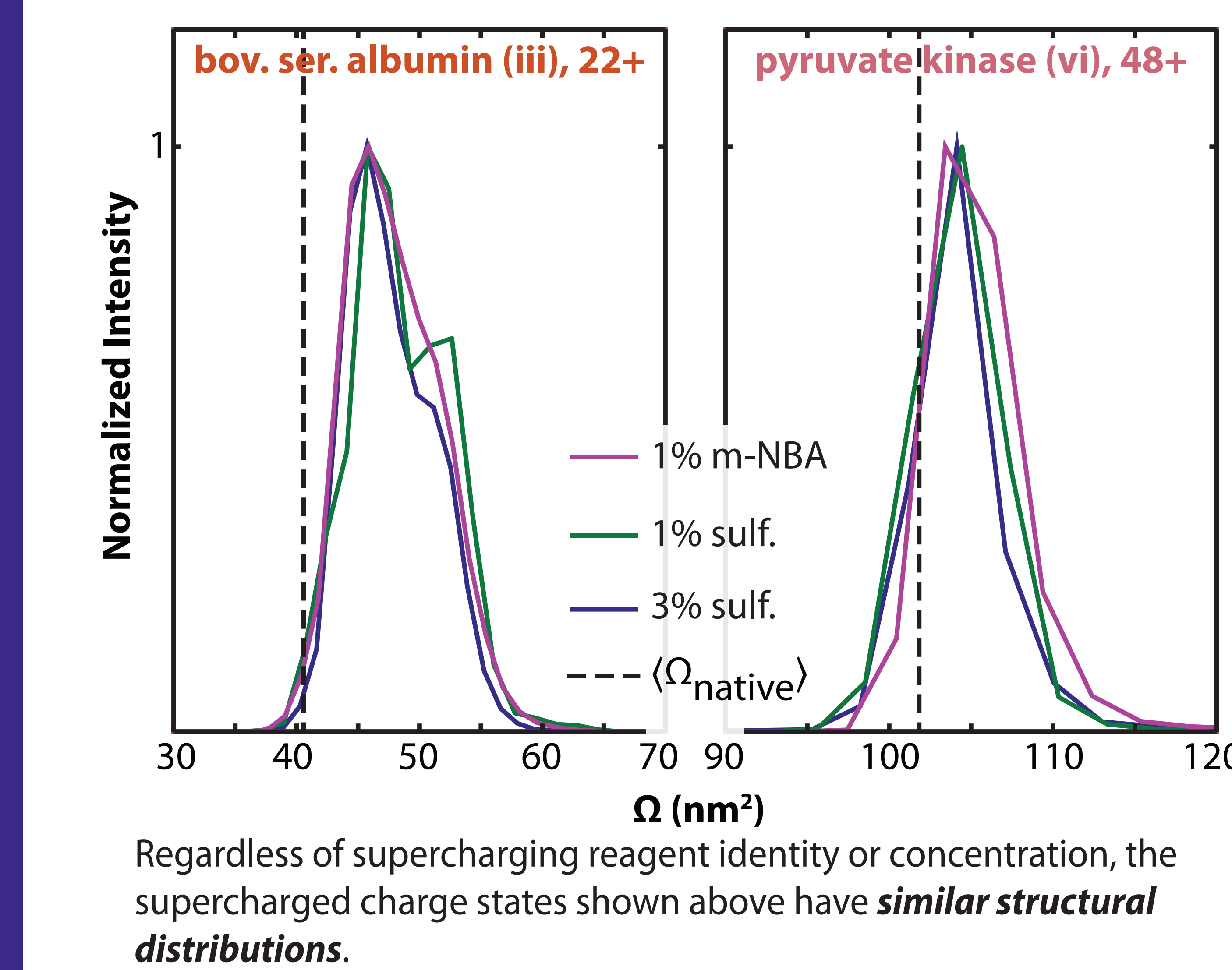
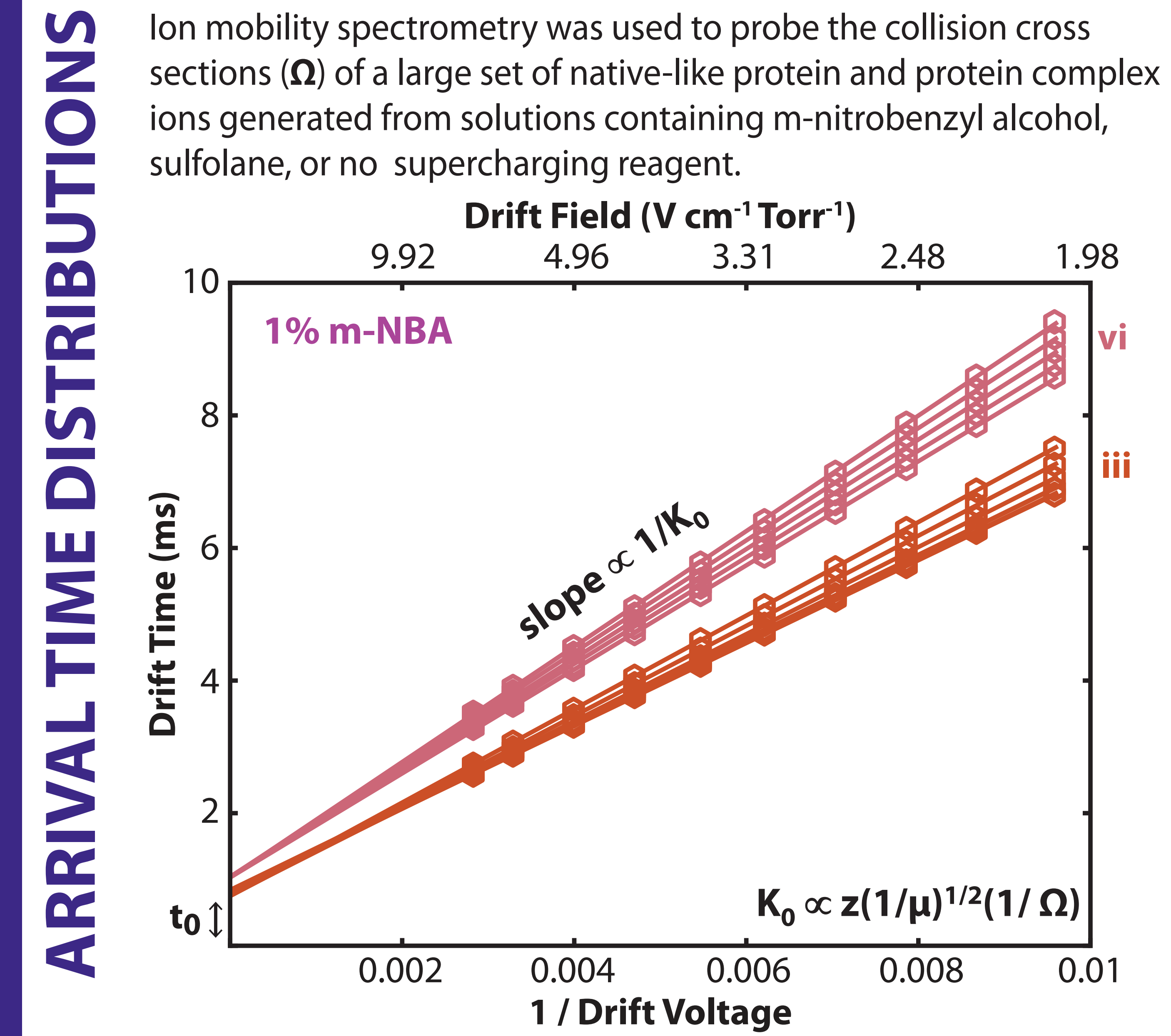
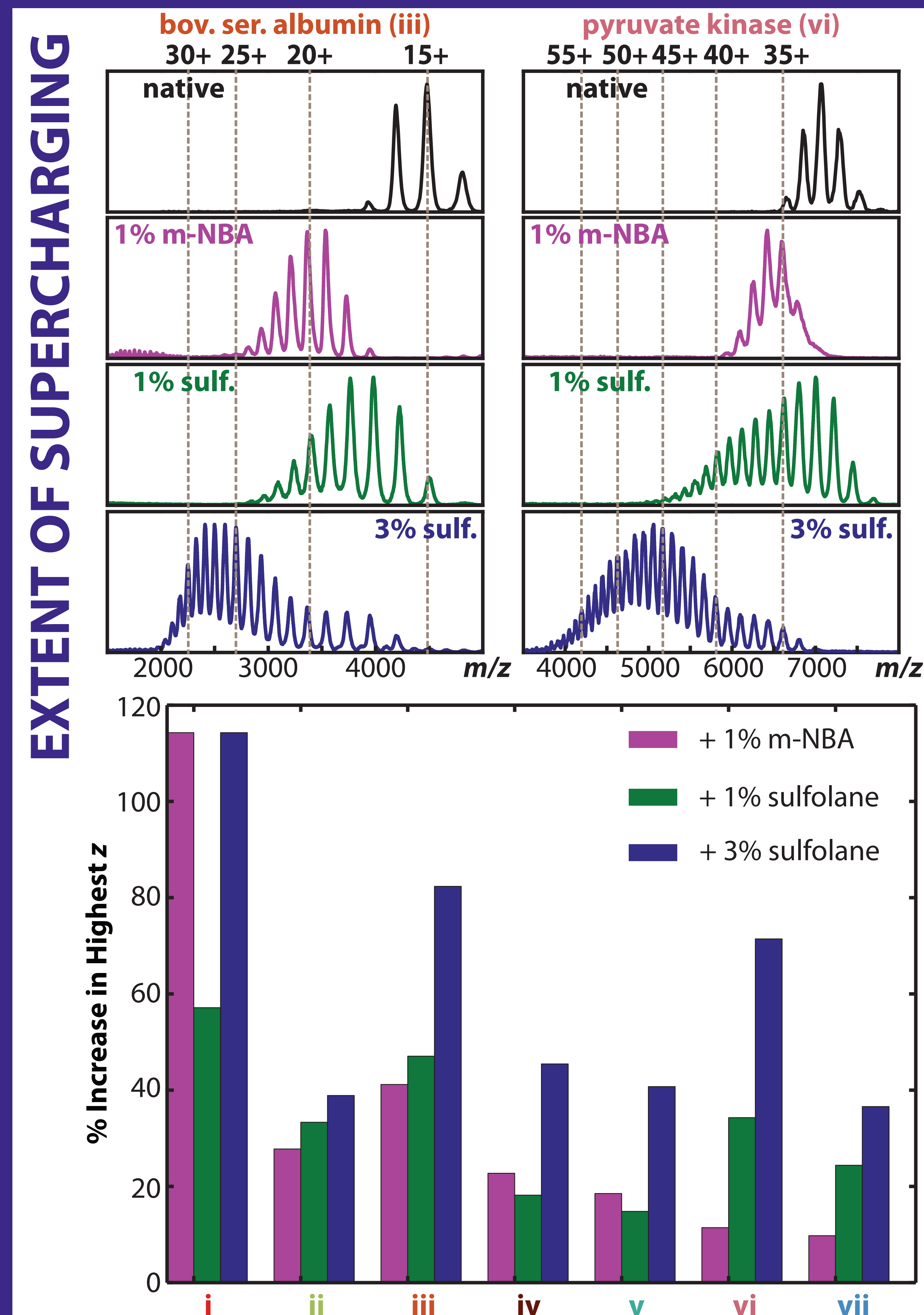
**Two Conflicting Theories:**

- 1) ...[supercharged charge states are concomitant with extended, non-native structures]...<sup>1,2</sup>
- 2) ...[supercharging does not affect native protein structure...any protein instability at high charge states is caused by the interactions of the protein with supercharging reagent]...<sup>3</sup>

Name	Mass (kDa)
i <b>cytochrome c</b>	12.4
ii <b>avidin</b>	64.0
iii <b>bovine serum albumin</b>	67.5
iv <b>concanavalin A</b>	103
v <b>alcohol dehydrogenase</b>	148
vi <b>pyruvate kinase</b>	233
vii <b>glutamate dehydrogenase</b>	337

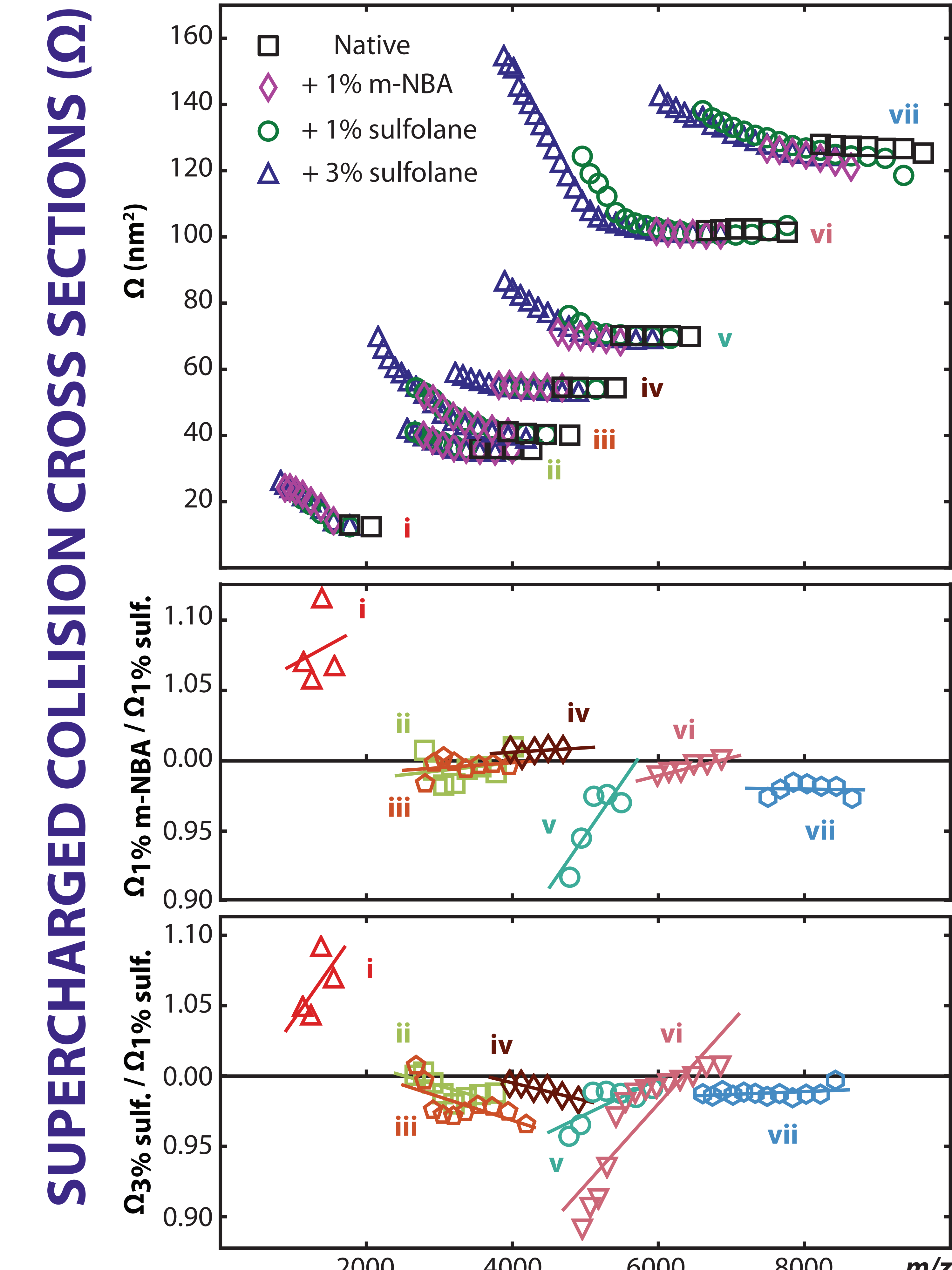
Unfolded Structure?

(1) Iavarone and Williams, *J. Am. Chem. Soc.*, **2003**, 125, 2319-2327.  
(2) Sterling, Daly, Feld, Thoren, Kintzer, Krantz and Williams, *J. Am. Soc. Mass Spectrom.*, **2010**, 21, 1762-1774.  
(3) Lomeli, Yin, Loo and Loo, *J. Am. Soc. Mass Spectrom.*, **2009**, 20, 593-596.



**CONCLUSIONS**

- A greater percent change in highest z was observed for smaller supercharged proteins (<64 kDa). Additionally, 1% m-NBA is a more effective supercharging reagent for smaller proteins, whereas 1-3% sulfolane is more effective for larger proteins (>64 kDa).
- Increases in  $\Omega$  with increasing charge state were much smaller for the largest protein complexes with and without supercharging.
- The results of this study suggest that the extent of supercharging on native protein structure depends strongly on reagent identity, but for most proteins and protein complexes, the  $\Omega$  of a given charge state depends weakly on supercharging reagent identity and concentration.
- Native protein structure can be retained with modest supercharging, however, the highest charge states exhibit varying degrees of unfolding.



The two plots above indicate that the  $\Omega$  values of proteins and protein complexes supercharged with either m-NBA or sulfolane are similar for a given charge state. We see that modest supercharging occurs with no change in  $\Omega$ , but that the highest charge states experience the greatest increase in  $\Omega$  and are thus associated with unfolding.

**THANKS**

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